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Leukaemia Section

Short Communication

t(11;19)(q23;p13) KMT2A/SH3GL1

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Abstract

Review on t(11;19)(q23;p13) KMT2A/SH3GL1, with data on clinics, and the genes implicated.

Identity

Note

At least three other translocation t(11;19) involve KMT2A: the t(11;19)(q23;p13.1) KMT2A/ELL, the t(11;19)(q23;p13.3) KMT2A/MLLT1, and the t(11;19)(q23;p13.3) KMT2A/ACER1.

Clinics and pathology

Epidemiology

Only one case to date, a 22-month-old girl with a diagnosis of acute myeloid leukemia not otherwise specified (So et al., 1997).

Genes involved and proteins

KMT2A

Location

11q23

Protein

KMT2A is better known as MLL. MLL is a transcriptional regulatory factor. MLL regulates chromatin-mediated transcription. MLL is a major regulator of hematopoiesis and embryonic development, through regulation of HOX genes expression. Translocations involving MLL are associated with high-risk de novo or treatment-related acute myeloid leukemias and

myelodysplastic syndromes and acute lymphoblastic leukemias (review in Huret, 2006). Mutations in MLL cause Wiedemann-Steiner syndrome (Jones et al., 2012).

SH3GL1

Location

19p13.3

Note

The gene has previously been named EEN.

Protein

The protein, named endophilin-A2, is made of 368 amino acids (aa) in isoform 1. From N-term to C-term, it contains a Bin/Amphiphysin/Rvs (BAR) domain (aa 18-249), involved in curvature of the cell membrane, a coiled-coil domain (protein-protein interactions, aa 119-227), and a SH3 domain (protein-protein interactions, aa 306-365). Endophilin-A2 is involved in clathrin-mediated endocytosis. Endophilin-A2 facilitates G1/S cell cycle transition of multiple myeloma cells (Huang et al., 2014).

Result of the chromosomal anomaly

Hybrid gene

Description

5' KMT2A - 3' SH3GL1. KMT2A exon 6 was fused to SH3GL1 exon 2.

Fusion protein

Description

1564 amino acids (1212 from KMT2A, and 352 from SH3GL1). The fusion protein retains the AT

hooks (DNA binding) and the Zinc finger CXXC-type from KMT2A fused to most of SH3GL1, including the BAR domain and the SH3 domain.

Oncogenesis

KMT2A/SH3GL1 is localized in the nucleus, while SH3GL1 is located in the cytoplasm. KMT2A/SH3GL1 may drive to aberrant transcriptional regulation. KMT2A/SH3GL1 increases HOXA7 promoter activity (Liu et al., 2004).

References

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